

21788wo.ST25.txt
SEQUENCE LISTING

<110> DSM IP Assets B.V.

<120> Feedback-resistant mevalonate kinases

<130> Case 21788

<150> EP 03012294.9

<151> 2003-06-12

<160> 33

<170> PatentIn version 3.2

<210> 1

<211> 378

<212> PRT

<213> Paracoccus zeaxanthinifaciens

<400> 1

Met Ser Thr Gly Arg Pro Glu Ala Gly Ala His Ala Pro Gly Lys Leu
1 5 10 15

Ile Leu Ser Gly Glu His Ser Val Leu Tyr Gly Ala Pro Ala Leu Ala
20 25 30

Met Ala Ile Ala Arg Tyr Thr Glu Val Trp Phe Thr Pro Leu Gly Ile
35 40 45

Gly Glu Gly Ile Arg Thr Thr Phe Ala Asn Leu Ser Gly Gly Ala Thr
50 55 60

Tyr Ser Leu Lys Leu Leu Ser Gly Phe Lys Ser Arg Leu Asp Arg Arg
65 70 75 80

Phe Glu Gln Phe Leu Asn Gly Asp Leu Lys Val His Lys Val Leu Thr
85 90 95

His Pro Asp Asp Leu Ala Val Tyr Ala Leu Ala Ser Leu Leu His Asp
100 105 110

Lys Pro Pro Gly Thr Ala Ala Met Pro Gly Ile Gly Ala Met His His
115 120 125

Leu Pro Arg Pro Gly Glu Leu Gly Ser Arg Thr Glu Leu Pro Ile Gly
130 135 140

Ala Gly Met Gly Ser Ser Ala Ala Ile Val Ala Ala Thr Thr Val Leu
145 150 155 160

Phe Glu Thr Leu Leu Asp Arg Pro Lys Thr Pro Glu Gln Arg Phe Asp
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170

175

Arg Val Arg Phe Cys Glu Arg Leu Lys His Gly Lys Ala Gly Pro Ile
 180 185 190

Asp Ala Ala Ser Val Val Arg Gly Gly Leu Val Arg Val Gly Gly Asn
 195 200 205

Gly Pro Gly Ser Ile Ser Ser Phe Asp Leu Pro Glu Asp His Asp Leu
 210 215 220

Val Ala Gly Arg Gly Trp Tyr Trp Val Leu His Gly Arg Pro Val Ser
 225 230 235 240

Gly Thr Gly Glu Cys Val Ser Ala Val Ala Ala Ala His Gly Arg Asp
 245 250 255

Ala Ala Leu Trp Asp Ala Phe Ala Val Cys Thr Arg Ala Leu Glu Ala
 260 265 270

Ala Leu Leu Ser Gly Gly Ser Pro Asp Ala Ala Ile Thr Glu Asn Gln
 275 280 285

Arg Leu Leu Glu Arg Ile Gly Val Val Pro Ala Ala Thr Gln Ala Leu
 290 295 300

Val Ala Gln Ile Glu Glu Ala Gly Gly Ala Ala Lys Ile Cys Gly Ala
 305 310 315 320

Gly Ser Val Arg Gly Asp His Gly Gly Ala Val Leu Val Arg Ile Asp
 325 330 335

Asp Ala Gln Ala Met Ala Ser Val Met Ala Arg His Pro Asp Leu Asp
 340 345 350

Trp Ala Pro Leu Arg Met Ser Arg Thr Gly Ala Ala Pro Gly Pro Ala
 355 360 365

Pro Arg Ala Gln Pro Leu Pro Gly Gln Gly
 370 375

<210> 2
 <211> 396
 <212> PRT
 <213> homo sapiens

<400> 2

Met Leu Ser Glu Val Leu Leu Val Ser Ala Pro Gly Lys Val Ile Leu
 Page 2

1	5	10	15
His Gly Glu His Ala Val Val His Gly Lys Val Ala Leu Ala Val Ser	20	25	30
Leu Asn Leu Arg Thr Phe Leu Arg Leu Gln Pro His Ser Asn Gly Lys	35	40	45
Val Asp Leu Ser Leu Pro Asn Ile Gly Ile Lys Arg Ala Trp Asp Val	50	55	60
Ala Arg Leu Gln Ser Leu Asp Thr Ser Phe Leu Glu Gln Gly Asp Val	65	70	75
Thr Thr Pro Thr Ser Glu Gln Val Glu Lys Leu Lys Glu Val Ala Gly	85	90	95
Leu Pro Asp Asp Cys Ala Val Thr Glu Arg Leu Ala Val Leu Ala Phe	100	105	110
Leu Tyr Leu Tyr Leu Ser Ile Cys Arg Lys Gln Arg Ala Leu Pro Ser	115	120	125
Leu Asp Ile Val Val Trp Ser Glu Leu Pro Pro Gly Ala Gly Leu Gly	130	135	140
Ser Ser Ala Ala Tyr Ser Val Cys Leu Ala Ala Ala Leu Leu Thr Val	145	150	155
Cys Glu Glu Ile Pro Asn Pro Leu Lys Asp Gly Asp Cys Val Asn Arg	165	170	175
Trp Thr Lys Glu Asp Leu Glu Leu Ile Asn Lys Trp Ala Phe Gln Gly	180	185	190
Glu Arg Met Ile His Gly Asn Pro Ser Gly Val Asp Asn Ala Val Ser	195	200	205
Thr Trp Gly Gly Ala Leu Arg Tyr His Gln Gly Lys Ile Ser Ser Leu	210	215	220
Lys Arg Ser Pro Ala Leu Gln Ile Leu Leu Thr Asn Thr Lys Val Pro	225	230	235
Arg Asn Thr Arg Ala Leu Val Ala Gly Val Arg Asn Arg Leu Leu Lys	245	250	255

Phe Pro Glu Ile Val Ala Pro Leu Leu Thr Ser Ile Asp Ala Ile Ser
 260 265 270

Leu Glu Cys Glu Arg Val Leu Gly Glu Met Gly Glu Ala Pro Ala Pro
 275 280 285

Glu Gln Tyr Leu Val Leu Glu Glu Leu Ile Asp Met Asn Gln His His
 290 295 300

Leu Asn Ala Leu Gly Val Gly His Ala Ser Leu Asp Gln Leu Cys Gln
 305 310 315 320

Val Thr Arg Ala Arg Gly Leu His Ser Lys Leu Thr Gly Ala Gly Gly
 325 330 335

Gly Gly Cys Gly Ile Thr Leu Leu Lys Pro Gly Leu Glu Gln Pro Glu
 340 345 350

Val Glu Ala Thr Lys Gln Ala Leu Thr Ser Cys Gly Phe Asp Cys Leu
 355 360 365

Glu Thr Ser Ile Gly Ala Pro Gly Val Ser Ile His Ser Ala Thr Ser
 370 375 380

Leu Asp Ser Arg Val Gln Gln Ala Leu Asp Gly Leu
 385 390 395

<210> 3
 <211> 395
 <212> PRT
 <213> mouse

<400> 3

Met Leu Ser Glu Ala Leu Leu Val Ser Ala Pro Gly Lys Val Ile Leu
 1 5 10 15

His Gly Glu His Ala Val Val His Gly Lys Val Ala Leu Ala Ala Ala
 20 25 30

Leu Asn Leu Arg Thr Phe Leu Leu Arg Pro Gln Ser Asn Gly Lys
 35 40 45

Val Ser Val Asn Leu Pro Asn Ile Gly Ile Lys Gln Val Trp Asp Val
 50 55 60

Gly Met Leu Gln Arg Leu Asp Thr Ser Phe Leu Glu Gln Gly Asp Val
 65 70 75 80

Ser Val Pro Thr Leu Glu Gln Leu Glu Lys Leu Lys Lys Met Gly Asp
 85 90 95
 Leu Pro Arg Asp Arg Ala Gly Asn Glu Gly Met Ala Leu Leu Ala Phe
 100 105 110
 Leu Tyr Leu Tyr Leu Ala Ile Cys Arg Lys Gln Arg Thr Leu Pro Ser
 115 120 125
 Leu Asp Met Val Val Trp Ser Glu Leu Pro Pro Gly Ala Gly Leu Gly
 130 135 140
 Ser Ser Ala Ala Tyr Ser Val Cys Leu Ala Ala Ala Leu Leu Thr Ala
 145 150 155 160
 Cys Glu Glu Val Ser Asn Pro Leu Lys Asp Gly Val Ser Val Ser Arg
 165 170 175
 Trp Pro Glu Glu Asp Leu Lys Ser Ile Asn Lys Trp Ala Phe Glu Gly
 180 185 190
 Glu Arg Val Ile His Gly Asn Pro Ser Gly Val Asp Asn Ala Val Ser
 195 200 205
 Thr Trp Gly Gly Ala Leu Arg Phe Gln Gln Gly Thr Met Ser Ser Leu
 210 215 220
 Lys Ser Leu Pro Ser Leu Gln Ile Leu Leu Thr Asn Thr Lys Val Pro
 225 230 235 240
 Arg Ser Thr Lys Ala Leu Val Ala Ala Val Arg Ser Arg Leu Thr Lys
 245 250 255
 Phe Pro Glu Ile Val Ala Pro Leu Leu Thr Ser Ile Asp Ala Ile Ser
 260 265 270
 Leu Glu Cys Glu Arg Val Leu Gly Glu Met Val Ala Ala Pro Val Pro
 275 280 285
 Glu Gln Tyr Leu Val Leu Glu Glu Leu Ile Asp Met Asn Gln His His
 290 295 300
 Leu Asn Ala Leu Gly Val Gly His Asn Ser Leu Asp Gln Leu Cys Gln
 305 310 315 320
 Val Thr Ala Ala His Gly Leu His Ser Lys Leu Thr Gly Ala Gly Gly
 325 330 335

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Gly Gly Cys Gly Ile Thr Leu Leu Lys Pro Gly Leu Glu Gln Ala Thr
340 345 350

Val Glu Ala Ala Lys Gln Ala Leu Thr Ser Cys Gly Phe Asp Cys Trp
355 360 365

Glu Thr Ser Ile Gly Ala Pro Gly Val Ser Thr His Ser Ala Ala Ala
370 375 380

Val Gly Asp Pro Val Arg Gln Ala Leu Gly Leu
385 390 395

<210> 4
<211> 395
<212> PRT
<213> rat

<400> 4

Met Leu Ser Glu Val Leu Leu Val Ser Ala Pro Gly Lys Val Ile Leu
1 5 10 15

His Gly Glu His Ala Val Val His Gly Lys Val Ala Leu Ala Val Ala
20 25 30

Leu Asn Leu Arg Thr Phe Leu Val Leu Arg Pro Gln Ser Asn Gly Lys
35 40 45

Val Ser Leu Asn Leu Pro Asn Val Gly Ile Lys Gln Val Trp Asp Val
50 55 60

Ala Thr Leu Gln Leu Leu Asp Thr Gly Phe Leu Glu Gln Gly Asp Val
65 70 75 80

Pro Ala Pro Thr Leu Glu Gln Leu Glu Lys Leu Lys Lys Val Ala Gly
85 90 95

Leu Pro Arg Asp Cys Val Gly Asn Glu Gly Leu Ser Leu Leu Ala Phe
100 105 110

Leu Tyr Leu Tyr Leu Ala Ile Cys Arg Lys Gln Arg Thr Leu Pro Ser
115 120 125

Leu Asp Ile Met Val Trp Ser Glu Leu Pro Pro Gly Ala Gly Leu Gly
130 135 140

Ser Ser Ala Ala Tyr Ser Val Cys Val Ala Ala Ala Leu Leu Thr Ala
145 150 155 160

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Cys Glu Glu Val Thr Asn Pro Leu Lys Asp Arg Gly Ser Ile Gly Ser
165 170 175

Trp Pro Glu Glu Asp Leu Lys Ser Ile Asn Lys Trp Ala Tyr Glu Gly
180 185 190

Glu Arg Val Ile His Gly Asn Pro Ser Gly Val Asp Asn Ser Val Ser
195 200 205

Thr Trp Gly Gly Ala Leu Arg Tyr Gln Gln Gly Lys Met Ser Ser Leu
210 215 220

Lys Arg Leu Pro Ala Leu Gln Ile Leu Leu Thr Asn Thr Lys Val Pro
225 230 235 240

Arg Ser Thr Lys Ala Leu Val Ala Gly Val Arg Ser Arg Leu Ile Lys
245 250 255

Phe Pro Glu Ile Met Ala Pro Leu Leu Thr Ser Ile Asp Ala Ile Ser
260 265 270

Leu Glu Cys Glu Arg Val Leu Gly Glu Met Ala Ala Ala Pro Val Pro
275 280 285

Glu Gln Tyr Leu Val Leu Glu Glu Leu Met Asp Met Asn Gln His His
290 295 300

Leu Asn Ala Leu Gly Val Gly His Ala Ser Leu Asp Gln Leu Cys Gln
305 310 315 320

Val Thr Ala Ala His Gly Leu His Ser Lys Leu Thr Gly Ala Gly Gly
325 330 335

Gly Gly Cys Gly Ile Thr Leu Leu Lys Pro Gly Leu Glu Arg Ala Lys
340 345 350

Val Glu Ala Ala Lys Gln Ala Leu Thr Gly Cys Gly Phe Asp Cys Trp
355 360 365

Glu Thr Ser Ile Gly Ala Pro Gly Val Ser Met His Ser Ala Thr Ser
370 375 380

Ile Glu Asp Pro Val Arg Gln Ala Leu Gly Leu
385 390 395

<210> 5
<211> 378

<212> PRT

<213> Arabidopsis thaliana

<400> 5

Met Glu Val Lys Ala Arg Ala Pro Gly Lys Ile Ile Leu Ala Gly Glu
 1 5 10 15

His Ala Val Val His Gly Ser Thr Ala Val Ala Ala Ala Ile Asp Leu
 20 25 30

Tyr Thr Tyr Val Thr Leu Arg Phe Pro Leu Pro Ser Ala Glu Asn Asn
 35 40 45

Asp Arg Leu Thr Leu Gln Leu Lys Asp Ile Ser Leu Glu Phe Ser Trp
 50 55 60

Ser Leu Ala Arg Ile Lys Glu Ala Ile Pro Tyr Asp Ser Ser Thr Leu
 65 70 75 80

Cys Arg Ser Thr Pro Ala Ser Cys Ser Glu Glu Thr Leu Lys Ser Ile
 85 90 95

Ala Val Leu Val Glu Glu Gln Asn Leu Pro Lys Glu Lys Met Trp Leu
 100 105 110

Ser Ser Gly Ile Ser Thr Phe Leu Trp Leu Tyr Thr Arg Ile Ile Gly
 115 120 125

Phe Asn Pro Ala Thr Val Val Ile Asn Ser Glu Leu Pro Tyr Gly Ser
 130 135 140

Gly Leu Gly Ser Ser Ala Ala Leu Cys Val Ala Leu Thr Ala Ala Leu
 145 150 155 160

Leu Ala Ser Ser Ile Ser Glu Lys Thr Arg Gly Asn Gly Trp Ser Ser
 165 170 175

Leu Asp Glu Thr Asn Leu Glu Leu Leu Asn Lys Trp Ala Phe Glu Gly
 180 185 190

Glu Lys Ile Ile His Gly Lys Pro Ser Gly Ile Asp Asn Thr Val Ser
 195 200 205

Ala Tyr Gly Asn Met Ile Lys Phe Cys Ser Gly Glu Ile Thr Arg Leu
 210 215 220

Gln Ser Asn Met Pro Leu Arg Met Leu Ile Thr Asn Thr Arg Val Gly
 225 230 235 240

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Arg Asn Thr Lys Ala Leu Val Ser Gly Val Ser Gln Arg Ala Val Arg
245 250 255

His Pro Asp Ala Met Lys Ser Val Phe Asn Ala Val Asp Ser Ile Ser
260 265 270

Lys Glu Leu Ala Ala Ile Ile Gln Ser Lys Asp Glu Thr Ser Val Thr
275 280 285

Glu Lys Glu Glu Arg Ile Lys Glu Leu Met Glu Met Asn Gln Gly Leu
290 295 300

Leu Leu Ser Met Gly Val Ser His Ser Ser Ile Glu Ala Val Ile Leu
305 310 315 320

Thr Thr Val Lys His Lys Leu Val Ser Lys Leu Thr Gly Ala Gly Gly
325 330 335

Gly Gly Cys Val Leu Thr Leu Leu Pro Thr Gly Thr Val Val Asp Lys
340 345 350

Val Val Glu Glu Leu Glu Ser Ser Gly Phe Gln Cys Phe Thr Ala Leu
355 360 365

Ile Gly Gly Asn Gly Ala Gln Ile Cys Tyr
370 375

<210> 6
<211> 443
<212> PRT
<213> Saccharomyces cerevisiae

<400> 6

Met Ser Leu Pro Phe Leu Thr Ser Ala Pro Gly Lys Val Ile Ile Phe
1 5 10 15

Gly Glu His Ser Ala Val Tyr Asn Lys Pro Ala Val Ala Ala Ser Val
20 25 30

Ser Ala Leu Arg Thr Tyr Leu Leu Ile Ser Glu Ser Ser Ala Pro Asp
35 40 45

Thr Ile Glu Leu Asp Phe Pro Asp Ile Ser Phe Asn His Lys Trp Ser
50 55 60

Ile Asn Asp Phe Asn Ala Ile Thr Glu Asp Gln Val Asn Ser Gln Lys
65 70 75 80

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Leu Ala Lys Ala Gln Gln Ala Thr Asp Gly Leu Ser Gln Glu Leu Val
 85 90 95
 Ser Leu Leu Asp Pro Leu Leu Ala Gln Leu Ser Glu Ser Phe His Tyr
 100 105 110
 His Ala Ala Phe Cys Phe Leu Tyr Met Phe Val Cys Leu Cys Pro His
 115 120 125
 Ala Lys Asn Ile Lys Phe Ser Leu Lys Ser Thr Leu Pro Ile Gly Ala
 130 135 140
 Gly Leu Gly Ser Ser Ala Ser Ile Ser Val Ser Leu Ala Leu Ala Met
 145 150 155 160
 Ala Tyr Leu Gly Gly Leu Ile Gly Ser Asn Asp Leu Glu Lys Leu Ser
 165 170 175
 Glu Asn Asp Lys His Ile Val Asn Gln Trp Ala Phe Ile Gly Glu Lys
 180 185 190
 Cys Ile His Gly Thr Pro Ser Gly Ile Asp Asn Ala Val Ala Thr Tyr
 195 200 205
 Gly Asn Ala Leu Leu Phe Glu Lys Asp Ser His Asn Gly Thr Ile Asn
 210 215 220
 Thr Asn Asn Phe Lys Phe Leu Asp Asp Phe Pro Ala Ile Pro Met Ile
 225 230 235 240
 Leu Thr Tyr Thr Arg Ile Pro Arg Ser Thr Lys Asp Leu Val Ala Arg
 245 250 255
 Val Arg Val Leu Val Thr Glu Lys Phe Pro Glu Val Met Lys Pro Ile
 260 265 270
 Leu Asp Ala Met Gly Glu Cys Ala Leu Gln Gly Leu Glu Ile Met Thr
 275 280 285
 Lys Leu Ser Lys Cys Lys Gly Thr Asp Asp Glu Ala Val Glu Thr Asn
 290 295 300
 Asn Glu Leu Tyr Glu Gln Leu Leu Glu Leu Ile Arg Ile Asn His Gly
 305 310 315 320
 Leu Leu Val Ser Ile Gly Val Ser His Pro Gly Leu Glu Leu Ile Lys

325

335

Asn Leu Ser Asp Asp Leu Arg Ile Gly Ser Thr Lys Leu Thr Gly Ala
340 345 350

Gly Gly Gly Gly Cys Ser Leu Thr Leu Leu Arg Arg Asp Ile Thr Gln
355 360 365

Glu Gln Ile Asp Ser Phe Lys Lys Lys Leu Gln Asp Asp Phe Ser Tyr
370 375 380

Glu Thr Phe Glu Thr Asp Leu Gly Gly Thr Gly Cys Cys Leu Leu Ser
385 390 395 400

Ala Lys Asn Leu Asn Lys Asp Leu Lys Ile Lys Ser Leu Val Phe Gln
405 410 415

Leu Phe Glu Asn Lys Thr Thr Thr Lys Gln Gln Ile Asp Asp Leu Leu
420 425 430

Leu Pro Gly Asn Thr Asn Leu Pro Trp Thr Ser
435 440

<210> 7
<211> 404
<212> PRT
<213> Schizosaccharomyces pombe

<400> 7

Met Ser Lys Ser Leu Ile Val Ser Ser Pro Gly Lys Thr Ile Leu Phe
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Gly Glu His Ala Val Val Tyr Gly Ala Thr Ala Leu Ala Ala Val
20 25 30

Ser Leu Arg Ser Tyr Cys Lys Leu Gln Thr Thr Asn Asn Asn Glu Ile
35 40 45

Val Ile Val Met Ser Asp Ile Gly Thr Glu Arg Arg Trp Asn Leu Gln
50 55 60

Ser Leu Pro Trp Gln His Val Thr Val Glu Asn Val Gln His Pro Ala
65 70 75 80

Ser Ser Pro Asn Leu Asp Leu Leu Gln Gly Leu Gly Glu Leu Leu Lys
85 90 95

Asn Glu Glu Asn Gly Leu Ile His Ser Ala Met Leu Cys Thr Leu Tyr
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Leu Phe Thr Ser Leu Ser Ser Pro Ser Gln Gly Cys Thr Leu Thr Ile
115 120 125

Ser Ser Gln Val Pro Leu Gly Ala Gly Leu Gly Ser Ser Ala Thr Ile
130 135 140

Ser Val Val Val Ala Thr Ser Leu Leu Leu Ala Phe Gly Asn Ile Glu
145 150 155 160

Pro Pro Ser Ser Asn Ser Leu Gln Asn Asn Lys Ala Leu Ala Leu Ile
165 170 175

Glu Ala Trp Ser Phe Leu Gly Glu Cys Cys Ile His Gly Thr Pro Ser
180 185 190

Gly Ile Asp Asn Ala Val Ala Thr Asn Gly Gly Leu Ile Ala Phe Arg
195 200 205

Lys Ala Thr Ala His Gln Ser Ala Met Lys Glu Phe Leu Lys Pro Lys
210 215 220

Asp Thr Leu Ser Val Met Ile Thr Asp Thr Lys Gln Pro Lys Ser Thr
225 230 235 240

Lys Lys Leu Val Gln Gly Val Phe Glu Leu Lys Glu Arg Leu Pro Thr
245 250 255

Val Ile Asp Ser Ile Ile Asp Ala Ile Asp Gly Ile Ser Lys Ser Ala
260 265 270

Val Leu Ala Leu Thr Ser Glu Ser Asp Lys Asn Ser Ser Ala Lys Lys
275 280 285

Leu Gly Glu Phe Ile Val Leu Asn Gln Lys Leu Leu Glu Cys Leu Gly
290 295 300

Val Ser His Tyr Ser Ile Asp Arg Val Leu Gln Ala Thr Lys Ser Ile
305 310 315 320

Gly Trp Thr Lys Leu Thr Gly Ala Gly Gly Gly Gly Cys Thr Ile Thr
325 330 335

Leu Leu Thr Pro Glu Cys Lys Glu Glu Glu Phe Lys Leu Cys Lys Glu
340 345 350

Ser Leu Leu Ala His Lys Asn Ser Ile Tyr Asp Val Gln Leu Gly Gly
 355 360 365

Pro Gly Val Ser Val Val Thr Asp Ser Asp Ser Phe Phe Pro Gln Tyr
 370 375 380

Glu Ser Asp Phe Asp Phe Lys Lys Leu Asn Leu Leu Ser Lys Phe Asn
 385 390 395 400

Lys Tyr Tyr Ile

<210> 8
 <211> 335
 <212> PRT
 <213> Pyrococcus abyssi

<400> 8

Met Pro Arg Leu Val Leu Ala Ser Ala Pro Ala Lys Ile Ile Leu Phe
 1 5 10 15

Gly Glu His Ser Val Val Tyr Gly Lys Pro Ala Ile Ala Ser Ala Ile
 20 25 30

Asp Leu Arg Thr Tyr Val Arg Ala Glu Phe Asn Asp Ser Gly Asn Ile
 35 40 45

Lys Ile Glu Ala His Asp Ile Lys Thr Pro Gly Leu Ile Val Ser Phe
 50 55 60

Ser Glu Asp Lys Ile Tyr Phe Glu Thr Asp Tyr Gly Lys Ala Ala Glu
 65 70 75 80

Val Leu Ser Tyr Val Arg His Ala Ile Glu Leu Val Leu Glu Glu Ala
 85 90 95

Asp Lys Arg Thr Gly Val Ser Val Ser Ile Thr Ser Gln Ile Pro Val
 100 105 110

Gly Ala Gly Leu Gly Ser Ser Ala Ala Val Ala Val Ala Thr Ile Gly
 115 120 125

Ala Val Ser Lys Leu Leu Asp Leu Glu Leu Ser Lys Glu Glu Ile Ala
 130 135 140

Lys Met Gly His Lys Val Glu Leu Leu Val Gln Gly Ala Ser Ser Gly
 145 150 155 160

Ile Asp Pro Thr Val Ser Ala Ile Gly Gly Phe Leu Tyr Tyr Lys Gln
165 170 175

Gly Glu Phe Glu His Leu Pro Phe Val Glu Leu Pro Ile Val Val Gly
180 185 190

Tyr Thr Gly Ser Ser Gly Ser Thr Lys Glu Leu Val Ala Met Val Arg
195 200 205

Arg Arg Tyr Glu Glu Met Pro Glu Leu Ile Glu Pro Ile Leu Glu Ser
210 215 220

Met Gly Lys Leu Val Asp Lys Ala Lys Glu Val Ile Ile Ser Lys Leu
225 230 235 240

Asp Glu Glu Glu Lys Phe Leu Lys Leu Gly Glu Leu Met Asn Ile Asn
245 250 255

His Gly Leu Leu Asp Ala Leu Gly Val Ser Thr Lys Lys Leu Ser Glu
260 265 270

Leu Val Tyr Ala Ala Arg Thr Ala Gly Ala Ile Gly Ala Lys Leu Thr
275 280 285

Gly Ala Gly Gly Gly Gly Cys Met Tyr Ala Leu Ala Pro Gly Lys Gln
290 295 300

Arg Glu Val Ala Thr Ala Ile Lys Ile Ala Gly Gly Thr Pro Met Ile
305 310 315 320

Thr Arg Ile Ser Lys Glu Gly Leu Arg Ile Glu Glu Val Arg Glu
325 330 335

<210> 9
<211> 335
<212> PRT
<213> Pyrococcus horikoshii

<400> 9

Met Val Lys Tyr Val Leu Ala Ser Ala Pro Ala Lys Val Ile Leu Phe
1 5 10 15

Gly Glu His Ser Val Val Tyr Gly Lys Pro Ala Ile Ala Ser Ala Ile
20 25 30

Glu Leu Arg Thr Tyr Val Arg Ala Gln Phe Asn Asp Ser Gly Asn Ile
35 40 45

Lys Ile Glu Ala His Asp Ile Lys Thr Pro Gly Leu Ile Val Ser Phe
 50 55 60
 Ser Glu Asp Lys Ile Tyr Phe Glu Thr Asp Tyr Gly Lys Ala Ala Glu
 65 70 75 80
 Val Leu Ser Tyr Val Arg Tyr Ala Ile Glu Leu Ala Leu Glu Glu Ser
 85 90 95
 Asp Lys Arg Val Gly Ile Asp Val Ser Ile Thr Ser Gln Ile Pro Val
 100 105 110
 Gly Ala Gly Leu Gly Ser Ser Ala Ala Val Ala Val Ala Thr Ile Gly
 115 120 125
 Ala Val Ser Arg Leu Leu Gly Leu Glu Leu Ser Lys Glu Glu Ile Ala
 130 135 140
 Lys Leu Gly His Lys Val Glu Leu Leu Val Gln Gly Ala Ser Ser Gly
 145 150 155 160
 Ile Asp Pro Thr Val Ser Ala Val Gly Gly Phe Leu Tyr Tyr Lys Gln
 165 170 175
 Gly Lys Phe Glu Pro Leu Pro Phe Met Glu Leu Pro Ile Val Val Gly
 180 185 190
 Tyr Thr Gly Ser Thr Gly Ser Thr Lys Glu Leu Val Ala Met Val Arg
 195 200 205
 Lys Arg Tyr Glu Glu Met Pro Glu Leu Val Glu Pro Ile Leu Glu Ala
 210 215 220
 Met Gly Lys Leu Val Asp Lys Ala Lys Glu Ile Ile Leu Ser Lys Leu
 225 230 235 240
 Asp Glu Glu Glu Lys Leu Thr Lys Leu Gly Glu Leu Met Asn Ile Asn
 245 250 255
 His Gly Leu Leu Asp Ala Leu Gly Val Ser Thr Lys Lys Leu Gly Glu
 260 265 270
 Leu Val Tyr Ala Ala Arg Thr Ala Gly Ala Ile Gly Ala Lys Leu Thr
 275 280 285
 Gly Ala Gly Gly Gly Gly Cys Met Tyr Ala Leu Ala Pro Gly Arg Gln
 290 295 300

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Arg Glu Val Ala Thr Ala Ile Lys Ile Ala Gly Gly Ile Pro Met Ile
305 310 315 320

Thr Arg Val Ser Arg Glu Gly Leu Arg Ile Glu Glu Val Ser Arg
325 330 335

<210> 10
<211> 334
<212> PRT
<213> Pyrococcus furiosus

<400> 10

Met Lys Val Ile Ala Ser Ala Pro Ala Lys Val Ile Leu Phe Gly Glu
1 5 10 15

His Ser Val Val Tyr Gly Lys Pro Ala Ile Ala Ala Ala Ile Asp Leu
20 25 30

Arg Thr Phe Val Glu Ala Glu Leu Ile Arg Glu Lys Lys Ile Arg Ile
35 40 45

Glu Ala His Asp Ile Lys Val Pro Gly Leu Thr Val Ser Phe Ser Glu
50 55 60

Asn Glu Ile Tyr Phe Glu Thr Asp Tyr Gly Lys Ala Ala Glu Val Leu
65 70 75 80

Ser Tyr Val Arg Glu Ala Ile Asn Leu Val Leu Glu Glu Ala Asp Lys
85 90 95

Lys Asn Val Gly Ile Lys Val Ser Ile Thr Ser Gln Ile Pro Val Gly
100 105 110

Ala Gly Leu Gly Ser Ser Ala Ala Val Ala Val Ala Thr Ile Gly Ala
115 120 125

Val Ser Lys Leu Leu Gly Leu Glu Leu Ser Lys Glu Glu Ile Ala Lys
130 135 140

Met Gly His Lys Thr Glu Leu Leu Val Gln Gly Ala Ser Ser Gly Ile
145 150 155 160

Asp Pro Thr Val Ser Ala Ile Gly Gly Phe Ile Phe Tyr Glu Lys Gly
165 170 175

Lys Phe Glu His Leu Pro Phe Met Glu Leu Pro Ile Val Val Gly Tyr
180 185 190

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Thr Gly Ser Ser Gly Pro Thr Lys Glu Leu Val Ala Met Val Arg Lys
195 200 205

Arg Tyr Glu Glu Met Pro Glu Leu Ile Val Pro Ile Leu Glu Ala Met
210 215 220

Gly Lys Val Val Glu Lys Ala Lys Asp Val Ile Leu Ser Asn Val Asp
225 230 235 240

Lys Glu Glu Lys Phe Glu Arg Leu Gly Val Leu Met Asn Ile Asn His
245 250 255

Gly Leu Leu Asp Ala Leu Gly Val Ser Thr Lys Lys Leu Ser Glu Leu
260 265 270

Val Tyr Ala Ala Arg Val Ala Gly Ala Leu Gly Ala Lys Ile Thr Gly
275 280 285

Ala Gly Gly Gly Gly Cys Met Tyr Ala Leu Ala Pro Asn Lys Gln Arg
290 295 300

Glu Val Ala Thr Ala Ile Arg Ile Ala Gly Gly Thr Pro Met Ile Thr
305 310 315 320

Glu Ile Ser Arg Glu Gly Leu Lys Ile Glu Glu Val Ile Lys
325 330

<210> 11
<211> 303
<212> PRT
<213> Methanobacterium thermoautotrophicum

<400> 11

Met Lys Ser Ser Ala Ser Ala Pro Ala Lys Ala Ile Leu Phe Gly Glu
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His Ala Val Val Tyr Ser Lys Pro Ala Ile Ala Ala Ala Ile Asp Arg
20 25 30

Arg Val Thr Val Thr Val Ser Glu Ser Ser Ser Thr His Val Thr Ile
35 40 45

Pro Ser Leu Gly Ile Arg His Ser Ser Glu Arg Pro Ser Gly Gly Ile
50 55 60

Leu Asp Tyr Ile Gly Arg Cys Leu Glu Leu Tyr His Asp Ala Ser Pro
65 70 75 80

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Leu Asp Ile Arg Val Glu Met Glu Ile Pro Ala Gly Ser Gly Leu Gly
85 90 95

Ser Ser Ala Ala Leu Thr Val Ala Leu Ile Gly Ala Leu Asp Arg Tyr
100 105 110

His Gly Arg Asp His Gly Pro Gly Glu Thr Ala Ala Arg Ala His Arg
115 120 125

Val Glu Val Asp Val Gln Gly Ala Ala Ser Pro Leu Asp Thr Ala Ile
130 135 140

Ser Thr Tyr Gly Gly Leu Val Tyr Leu Asp Ser Gln Arg Arg Val Arg
145 150 155 160

Gln Phe Glu Ala Asp Leu Gly Asp Leu Val Ile Ala His Leu Asp Tyr
165 170 175

Ser Gly Glu Thr Ala Arg Met Val Ala Gly Val Ala Glu Arg Phe Arg
180 185 190

Arg Phe Pro Asp Ile Met Gly Arg Ile Met Asp Thr Val Glu Ser Ile
195 200 205

Thr Asn Thr Ala Tyr Arg Glu Leu Leu Arg Asn Asn Thr Glu Pro Leu
210 215 220

Gly Glu Leu Met Asn Leu Asn Gln Gly Leu Leu Asp Ser Met Gly Val
225 230 235 240

Ser Thr Arg Glu Leu Ser Met Met Val Tyr Glu Ala Arg Asn Ala Gly
245 250 255

Ala Ala Gly Ser Lys Ile Thr Gly Ala Gly Gly Gly Gly Ser Ile Ile
260 265 270

Ala His Cys Pro Gly Cys Val Asp Asp Val Val Thr Ala Leu Asn Arg
275 280 285

Asn Trp Lys Ala Met Arg Ala Glu Phe Ser Val Lys Gly Leu Ile
290 295 300

<210> 12
<211> 284
<212> PRT
<213> Archaeoglobus fulgidus
<400> 12

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Met Ile Ala Ser Ala Pro Gly Lys Ile Ile Leu Phe Gly Glu His Ala
1 5 10 15

Val Val Tyr Gly Arg His Ala Val Val Ser Ala Ile Asn Leu Arg Cys
20 25 30

Arg Val Ser Val Arg Lys Ser Asp Arg Phe Leu Ile Arg Ser Ser Leu
35 40 45

Gly Glu Ser Gly Leu Asp Tyr Gln Arg His Pro Tyr Val Val Gln Ala
50 55 60

Val Lys Arg Phe Gly Glu Leu Arg Asn Ile Pro Gly Ala Glu Ile Glu
65 70 75 80

Ile Glu Ser Glu Ile Pro Ile Gly Ser Gly Leu Gly Ser Ser Ala Ala
85 90 95

Val Ile Val Ala Thr Ile Ala Ala Leu Asn Ala Glu Phe Asp Gly Asp
100 105 110

Met Asp Lys Glu Ala Ile Phe Gln Met Ala Lys Gln Val Glu Ile Asp
115 120 125

Val Gln Gly Arg Ala Ser Gly Ile Asp Pro Phe Ile Ser Thr Phe Gly
130 135 140

Gly Ser Trp Leu Phe Pro Glu Arg Arg Lys Val Glu Met Pro Phe Lys
145 150 155 160

Phe Phe Val Ile Asn Phe Gly Ser Arg Ser Thr Ala Glu Met Val Ala
165 170 175

Lys Val Ala Glu Leu Arg Glu Arg His Pro Glu Val Val Asp Lys Ile
180 185 190

Phe Asp Ala Ile Asp Ala Ile Ser Leu Glu Ala Ser Asp Val Gly Ser
195 200 205

Ala Glu Arg Leu Glu Glu Leu Ile Ala Ile Asn Gln Ser Leu Leu Arg
210 215 220

Ala Ile Gly Val Ser Asn Pro Glu Ile Asp Arg Thr Ile Ala Glu Leu
225 230 235 240

Glu Arg Met Gly Leu Asn Ala Lys Ile Thr Gly Ala Gly Gly Gly Gly
245 250 255

21788wo.ST25.txt

Cys Ile Phe Gly Leu Phe Lys Gly Glu Lys Pro Lys Gly Ser Phe Ile
260 265 270

Val Glu Pro Glu Lys Glu Gly Val Arg Ile Glu Glu
275 280

<210> 13
<211> 312
<212> PRT
<213> Methanococcus jannaschii

<400> 13

Met Ile Ile Glu Thr Pro Ser Lys Val Ile Leu Phe Gly Glu His Ala
1 5 10 15

Val Val Tyr Gly Tyr Arg Ala Ile Ser Met Ala Ile Asp Leu Thr Ser
20 25 30

Thr Ile Glu Ile Lys Glu Thr Gln Glu Asp Glu Ile Ile Leu Asn Leu
35 40 45

Asn Asp Leu Asn Lys Ser Leu Gly Leu Asn Leu Asn Glu Ile Lys Asn
50 55 60

Ile Asn Pro Asn Asn Phe Gly Asp Phe Lys Tyr Cys Leu Cys Ala Ile
65 70 75 80

Lys Asn Thr Leu Asp Tyr Leu Asn Ile Glu Pro Lys Thr Gly Phe Lys
85 90 95

Ile Asn Ile Ser Ser Lys Ile Pro Ile Ser Cys Gly Leu Gly Ser Ser
100 105 110

Ala Ser Ile Thr Ile Gly Thr Ile Lys Ala Val Ser Gly Phe Tyr Asn
115 120 125

Lys Glu Leu Lys Asp Asp Glu Ile Ala Lys Leu Gly Tyr Met Val Glu
130 135 140

Lys Glu Ile Gln Gly Lys Ala Ser Ile Thr Asp Thr Ser Thr Ile Thr
145 150 155 160

Tyr Lys Gly Ile Leu Glu Ile Lys Asn Asn Lys Phe Arg Lys Ile Lys
165 170 175

Gly Glu Phe Glu Glu Phe Leu Lys Asn Cys Lys Phe Leu Ile Val Tyr
180 185 190

21788wo.ST25.txt

Ala Glu Lys Arg Lys Lys Lys Thr Ala Glu Leu Val Asn Glu Val Ala
195 200 205

Lys Ile Glu Asn Lys Asp Glu Ile Phe Lys Glu Ile Asp Lys Val Ile
210 215 220

Asp Glu Ala Leu Lys Ile Lys Asn Lys Glu Asp Phe Gly Lys Leu Met
225 230 235 240

Thr Lys Asn His Glu Leu Leu Lys Lys Leu Asn Ile Ser Thr Pro Lys
245 250 255

Leu Asp Arg Ile Val Asp Ile Gly Asn Arg Phe Gly Phe Gly Ala Lys
260 265 270

Leu Thr Gly Ala Gly Gly Gly Gly Cys Val Ile Ile Leu Val Asn Glu
275 280 285

Glu Lys Glu Lys Glu Leu Leu Lys Glu Leu Asn Lys Glu Asp Val Arg
290 295 300

Ile Phe Asn Cys Arg Met Met Asn
305 310

<210> 14
<211> 324
<212> PRT
<213> Aeropyrum pernix

<400> 14

Met Arg Arg Ala Ala Arg Ala Ser Ala Pro Gly Lys Val Ile Ile Val
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Gly Glu His Phe Val Val Arg Gly Ser Leu Ala Ile Val Ala Ala Ile
20 25 30

Gly Arg Arg Leu Arg Val Thr Val Arg Ser Gly Gly Lys Gly Ile Val
35 40 45

Leu Glu Ser Ser Met Leu Gly Arg His Ser Ala Pro Leu Pro Gly Gln
50 55 60

Gly Ala Ala Ala Lys Val Ser Pro Val Leu Glu Pro Tyr Ile Ala Val
65 70 75 80

Leu Arg Ser Leu Ala Ala Arg Gly Tyr Ser Val Val Pro His Thr Ile
85 90 95

21788wo.ST25.txt

Leu Val Glu Ser Gly Ile Pro Pro Arg Ala Gly Leu Gly Ser Ser Ala
 100 105 110
 Ala Ser Met Val Ala Tyr Ala Leu Ser Tyr Ser Ala Met His Gly Asp
 115 120 125
 Pro Leu Ser Ala Glu Asp Leu Tyr Ser Val Ala Met Glu Gly Glu Lys
 130 135 140
 Ile Ala His Gly Lys Pro Ser Gly Val Asp Val Thr Ile Ala Val Arg
 145 150 155 160
 Gly Gly Val Leu Ala Tyr Arg Arg Gly Glu Asn Pro Val Asp Ile Arg
 165 170 175
 Pro Gly Leu Thr Gly Val Thr Leu Leu Val Ala Asp Thr Gly Val Glu
 180 185 190
 Arg Arg Thr Arg Asp Val Val Glu His Val Leu Ser Ile Ala Asp Ala
 195 200 205
 Leu Gly Glu Ala Ser Thr Tyr Ile Tyr Arg Ala Ala Asp Leu Ile Ala
 210 215 220
 Arg Glu Ala Leu His Ala Ile Glu Lys Gly Asp Ala Glu Arg Leu Gly
 225 230 235 240
 Leu Ile Met Asn Ala Ala Gln Gly Leu Leu Ser Ser Leu Gly Ala Ser
 245 250 255
 Ser Leu Glu Ile Glu Thr Leu Val Tyr Arg Met Arg Ser Ala Gly Ala
 260 265 270
 Leu Gly Ala Lys Leu Thr Gly Ala Gly Trp Gly Gly Cys Val Ile Gly
 275 280 285
 Leu Phe Lys Glu Gly Glu Val Glu Arg Gly Leu Glu Ser Val Val Glu
 290 295 300
 Ser Ser Ser Gln Ala Phe Thr Ala Ser Ile Ala Glu Glu Gly Ala Arg
 305 310 315 320
 Leu Glu Glu Phe

<211> 387
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> His6-Paracoccus zeaxanthinifaciens

<400> 15

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 1 5 10 15

Ala Gly Ala His Ala Pro Gly Lys Leu Ile Leu Ser Gly Glu His Ser
 20 25 30

Val Leu Tyr Gly Ala Pro Ala Leu Ala Met Ala Ile Ala Arg Tyr Thr
 35 40 45

Glu Val Trp Phe Thr Pro Leu Gly Ile Gly Glu Gly Ile Arg Thr Thr
 50 55 60

Phe Ala Asn Leu Ser Gly Gly Ala Thr Tyr Ser Leu Lys Leu Leu Ser
 65 70 75 80

Gly Phe Lys Ser Arg Leu Asp Arg Arg Phe Glu Gln Phe Leu Asn Gly
 85 90 95

Asp Leu Lys Val His Lys Val Leu Thr His Pro Asp Asp Leu Ala Val
 100 105 110

Tyr Ala Leu Ala Ser Leu Leu His Asp Lys Pro Pro Gly Thr Ala Ala
 115 120 125

Met Pro Gly Ile Gly Ala Met His His Leu Pro Arg Pro Gly Glu Leu
 130 135 140

Gly Ser Arg Thr Glu Leu Pro Ile Gly Ala Gly Met Gly Ser Ser Ala
 145 150 155 160

Ala Ile Val Ala Ala Thr Thr Val Leu Phe Glu Thr Leu Leu Asp Arg
 165 170 175

Pro Lys Thr Pro Glu Gln Arg Phe Asp Arg Val Arg Phe Cys Glu Arg
 180 185 190

Leu Lys His Gly Lys Ala Gly Pro Ile Asp Ala Ala Ser Val Val Arg
 195 200 205

Gly Gly Leu Val Arg Val Gly Gly Asn Gly Pro Gly Ser Ile Ser Ser
 210 215 220

21788wo.ST25.txt

Phe Asp Leu Pro Glu Asp His Asp Leu Val Ala Gly Arg Gly Trp Tyr
225 230 235 240

Trp Val Leu His Gly Arg Pro Val Ser Gly Thr Gly Glu Cys Val Ser
245 250 255

Ala Val Ala Ala Ala His Gly Arg Asp Ala Ala Leu Trp Asp Ala Phe
260 265 270

Ala Val Cys Thr Arg Ala Leu Glu Ala Ala Leu Leu Ser Gly Gly Ser
275 280 285

Pro Asp Ala Ala Ile Thr Glu Asn Gln Arg Leu Leu Glu Arg Ile Gly
290 295 300

Val Val Pro Ala Ala Thr Gln Ala Leu Val Ala Gln Ile Glu Glu Ala
305 310 315 320

Gly Gly Ala Ala Lys Ile Cys Gly Ala Gly Ser Val Arg Gly Asp His
325 330 335

Gly Gly Ala Val Leu Val Arg Ile Asp Asp Ala Gln Ala Met Ala Ser
340 345 350

Val Met Ala Arg His Pro Asp Leu Asp Trp Ala Pro Leu Arg Met Ser
355 360 365

Arg Thr Gly Ala Ala Pro Gly Pro Ala Pro Arg Ala Gln Pro Leu Pro
370 375 380

Gly Gln Gly
385

<210> 16
<211> 1137
<212> DNA
<213> Paracoccus zeaxanthinifaciens

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gtgtggttca cgccgcttgg cattggcgag gggatacgca cgacattcgc caatctctcg 180
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ttcgagcagt tcctgaacgg cgacctaaag gtgcacaagg tcctgaccca tcccgcgat 300
ctggcggtct atgcgctggc gtcgcttctg cacgacaagc cgccggggac cgccgcgatg 360

21788wo.ST25.txt

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gggaccggcg	aatgcgtcag	cgcggtcgcg	gcggcgcatg	gtcgcgatgc	ggcgctgtgg	780
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atggcttcgg	tcattggcgcg	ccatcccagc	ctcgactggg	cgcccctgcg	catgtcgcgc	1080
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<210> 17
 <211> 1191
 <212> DNA
 <213> homo sapiens

<400> 17						
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cttcaacccc	acagcaatgg	gaaagtggac	ctcagcttac	ccaacattgg	tatcaagcgg	180
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acaacaccca	cctcagagca	agtggagaag	ctaaaggagg	ttgcaggctt	gcctgacgac	300
tgtgctgtca	ccgagcgcct	ggctgtgctg	gcctttcttt	acttatacct	gtccatctgc	360
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cgcaatacca	gggcccttgt	ggctggcgctc	agaaacaggc	tgctcaagtt	cccagagatc	780
gtggcccccc	tcctgacctc	aatagatgcc	atctccctgg	agtgtgagcg	cgtgctggga	840
gagatggggg	aagccccagc	cccggagcag	tacctcgtgc	tggaagagct	cattgacatg	900

21788wo.ST25.txt

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aaccagcacc atctgaatgc cctcggcgtg ggccacgcct ctctggacca gctctgccag 960
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atcacactcc tcaagccagg gctggagcag ccagaagtgg aggccacgaa gcaggccctg 1080
accagctgtg gctttgactg cttggaaacc agcatcgggtg cccccggcgt ctccatccac 1140
tcagccacct ccctggacag ccgagtccag caagccctgg atggcctctg a 1191

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```

<210> 18
<211> 1188
<212> DNA
<213> mouse

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<400> 18
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ctgcgaccgc agagcaatgg gaaagtgagc gtcaatttac ccaacatcgg tattaagcag 180
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cgtgcaggca atgaaggcat ggctctgctt gcctttctct acctgtacct ggcaatctgc 360
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tcagctgcag ctgtagggga ccctgtccga caagccctgg gcctctga 1188

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<210> 19
<211> 1188
<212> DNA
<213> rat

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21788wo.ST25.txt

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<400> 19
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ctgcgaccgc agagcaatgg gaaagtgagc ctcaatttac caaacgtcgg tattaagcag    180
gtctgggatg tggccacact tcagctgctg gacacaggct ttcttgagca aggcgatgtc    240
ccggcaccca ccttgagca actggagaag ctgaagaagg tggcgggcct ccccgagac    300
tgtgtaggca acgaaggcct gtctctgctt gcctttctgt acctgtacct ggctatctgc    360
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gcgggcttgg gctccagtgc agcctactcg gtgtgtgtgg cagccgccct cctgactgcc    480
tgtgaggagg tcaccaaccc gctcaaggac aggggctcca ttggcagttg gcccaggagg    540
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aaccagcacc atctgaatgc ccttggtgtg ggccacgcct ccctggacca gctctgtcag    960
gtaacagcag cacatggact gcacagcaag ctgactggcg caggcggcgg cggctgtggc   1020
atcacctcc tgaagccagg tctagagcga gcaaaagtgg aggccgcaa gcaggccctg   1080
accggctgcg ggtttgactg ctgggagacc agcattggag cgctgggggt ctccatgcac   1140
tcagccacct ccatagagga ccctgtccga caagccctgg gcctctga               1188

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<210> 20
<211> 1137
<212> DNA
<213> Arabidopsis thaliana

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<400> 20
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cctcttccat cagctgagaa caatgatagg cttacacttc agctcaagga catttccttg    180
gagttttcat ggtccttagc cagaatcaaa gaagcgattc cttatgattc aagcactctc    240
tgccgttcta cgccggcttc atgttcagag gagaccctta aatcaattgc agttttgggt    300
gaagagcaaa atcttccaaa ggaaaagatg tggctctcct ctgggatctc cacgtttctc    360
tggttataca ccagaattat agggttcaat ccggctacag tagtcattaa ctctgagctt    420

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21788wo.ST25.txt

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ataactcggg tacaatccaa catgcctctg agaatgctaa ttaccaacac tagagttggg	720
cgaaacacaa aagctctggg ctctgggtgtg tcacagagag cggtaaagaca tcctgatgcg	780
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tctaaagacg agacctcagt tacagaaaaa gaagagagaa taaaagaact catggagatg	900
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accacgggtca agcacaagct tgtctccaaa cttacaggag ctggtggcgg cggctgcgtc	1020
ctcactctat taccâaccgg gacgggtggg gacaaagtgg tggaggagct cgagtcagc	1080
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<210> 21
 <211> 1332
 <212> DNA
 <213> *Saccharomyces cerevisiae*

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ccgttggttag ctcaactatc cgaatccttc cactaccatg cagcgttttg tttcctgtat	360
atgtttgttt gcctatgccc ccatgccaaag aatattaagt tttctttaaa gtctacttta	420
cccatcgggtg ctgggttggg ctcaagcgcc tctatttctg tatcactggc cttagctatg	480
gcctacttgg ggggggttaat aggatctaag gacttggaag agctgtcaga aaacgataag	540
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21788wo.ST25.txt

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gcaaaaaatt tgaataaaga tcttaaaatc aaatccctag tattccaatt atttgaaaat	1260
aaaactacca caaagcaaca aattgacgat ctattattgc caggaaacac gaatttacca	1320
tggacttcat aa	1332

<210> 22

<211> 1215

<212> DNA

<213> Schizosaccharomyces pombe

<400> 22

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cagacgacta ataacaatga aatagtaatt gtgatgagtg atatagggac cgaacgccga	180
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21788wo.ST25.txt

<210> 23
<211> 1008
<212> DNA
<213> Pyrococcus abyssi

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cacctaccat tcgtggagct tccaatagta gttggatata ccggctcaag tggctccaca 600
aaggaattag ttgcatggt taggagaagg tacgaggaga tgcccagatt aattgaaccc 660
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<210> 24
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<212> DNA
<213> Pyrococcus horikoshii

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